

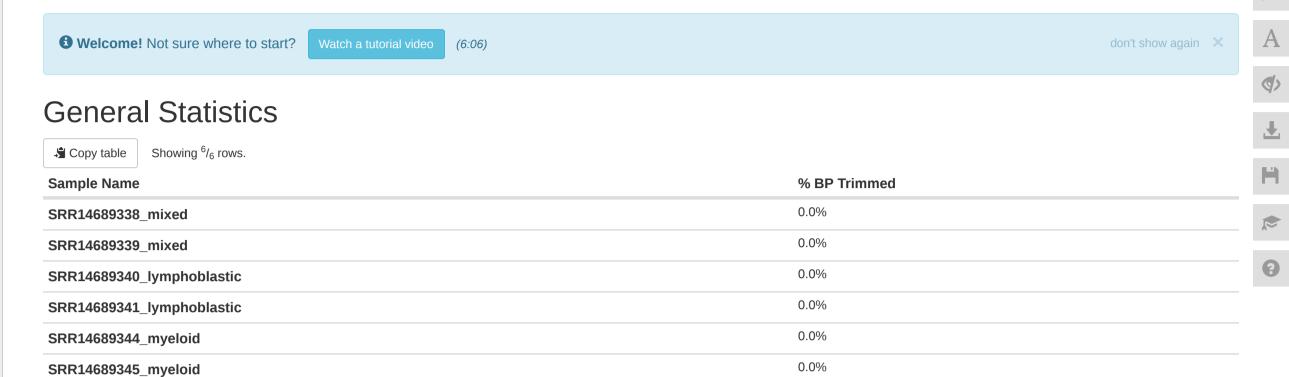
General Stats

Cutadapt



A modular tool to aggregate results from bioinformatics analyses across many samples into a single report.

Report generated on 2022-04-23, 09:03 based on data in: /home/khoidnyds/RNAseq_old/3.cutadapt



Cutadapt

Cutadapt is a tool to find and remove adapter sequences, primers, poly-A tails and other types of unwanted sequence from your high-throughput sequencing reads. DOI: 10.14806/ej.17.1.200.

Filtered Reads

This plot shows the number of reads (SE) / pairs (PE) removed by Cutadapt.

